[0135] The recombinant cloning vector, according to this disclosure, then comprises the selected DNA of the DNA sequences of this disclosure for expression in a suitable host. The DNA is operatively linked in the vector to an expression control sequence in the recombinant DNA molecule so that the actin polypeptide can be expressed. The expression control sequence may be selected from the group consisting of sequences that control the expression of genes of prokaryotic or eukaryotic cells and their viruses and combinations thereof. The expression control sequence may be specifically selected from the group consisting of the lac system, the trp system, the tac system, the trc system, major operator and promoter regions of phage lambda, the control region of fd coat protein, the early and late promoters of SV40, promoters derived from polyoma, adenovirus, retrovirus, baculovirus and simian virus, the promoter for 3-phosphoglycerate kinase, the promoters of yeast acid phosphatase, the promoter of the yeast alpha-mating factors and combinations thereof.

EXAMPLE 5

Motor Protein Variants

[0136] Variants of the motor proteins (such as actin and myosin) can be used instead of the native proteins, as long as the variants retain the motor activity. DNA mutagenesis techniques may be used to produce variant DNA molecules, and will facilitate the production of proteins which differ in certain structural aspects from the native protein, yet the variant proteins are clearly derivative and maintain the essential functional characteristic of the motor protein as defined above. Newly derived proteins may also be selected in order to obtain variations in the characteristics of the motor protein, as will be more fully described below. Such derivatives include those with variations in the amino acid sequence including minor deletions, additions and substitutions.

[0137] While the site for introducing an amino acid sequence variation is predetermined, the mutation per se need not be predetermined. For example, in order to optimize the performance of a mutation at a given site, random mutagenesis may be conducted at a target codon or region and the expressed protein variants screened for optimal activity. Techniques for making substitution mutations at predetermined sites in DNA having a known sequence are well known.

[0138] Amino acid substitutions are typically of single residues, for example 1, 2, 3, 4 or more substitutions; insertions usually will be on the order of about from 1 to 10 amino acid residues; and deletions will range about from 1 to 30 residues. Substitutions, deletions, insertions or any combination thereof may be combined to arrive at a final construct. Obviously, the mutations that are made in the DNA encoding the protein must not place the sequence out of reading frame, and preferably will not create complementary regions that could produce secondary changes in the mRNA structure.

[0139] Substitutional variants are those in which at least one residue in the amino acid sequence has been removed and a different residue inserted in its place. Such substitutions are generally conservative substitutions when it is desired to finely modulate the characteristics of the protein.

Examples of such conservative substitutions are well known, and are shown, for example, in U.S. Pat. No. 5,928,896 and U.S. Pat. No. 5,917,019.

[0140] Substantial changes in function or immunological identity are made by selecting substitutions that are less conservative i.e., selecting residues that differ more significantly in their effect on maintaining (a) the structure of the polypeptide backbone in the area of the substitution, for example, as a sheet or helical conformation, (b) the charge or hydrophobicity of the molecule at the target site, or (c) the bulk of the side chain. The substitutions which in general are expected to produce the greatest changes in protein properties will be those in which (a) a hydrophilic residue, e.g., seryl or threonyl, is substituted for (or by) a hydrophobic residue, e.g., leucyl, isoleucyl, phenylalanyl, valyl or alanyl; (b) a cysteine or proline is substituted for (or by) any other residue; (c) a residue having an electropositive side chain, e.g., lysyl, arginyl, or histadyl, is substituted for (or by) an electronegative residue, e.g., glutamyl or aspartyl; or (d) a residue having a bulky side chain, e.g., phenylalanine, is substituted for (or by) one not having a side chain, e.g., glycine.

EXAMPLE 6

[0141] An embodiment of a molecular motor 310 that includes annular substrates is depicted in FIG. 10. Discs are shown as the annular substrates in FIG. 10, but a layer of concentric rings lying in a common plane may be substituted for each one of the discs. These rings and ring layers are shown in detail in FIG. 13.

[0142] With reference to FIG. 10, a planar surface of a first disc 311 is secured to a base 312 so that the first disc 311 is not free to rotate relative to the base 312. The first disc 311 may be secured to the base 312 by any suitable manner such as by an adhesive. A second disc 313 is secured to a drive member 314 so that the second disc 313 is free to rotate relative to the first disc 311. The second disc 313 may be secured to the drive member 314 by any suitable manner such as by an adhesive. The drive member 314 may include a series of gear teeth for driving a driven member similar to that shown in FIG. 1. The first disc 311 and the second disc 313 are axially aligned relative to each other along a central longitudinal axis 320. The first disc 311 and the second disc 313 each define a respective orifice (depicted, for example, as element 352 in FIG. 12A or as element 372 in FIG. 12B) centered on the central axis 320. The orifices receive a support rod 319 that is axially aligned along the central axis 320. The support rod 319 is secured by the base 312 so that the support rod 319 is not free to rotate relative to the base 312. The support rod 319 is received within the drive member 314 so that the drive member 314 and second disc 313 remain free to rotate relative to the support rod 319. Bushings or ball bearings (not shown) may be provided at the surface interfaces between the support rod 319 and the drive member 314, and between the support rod 319 and the second disc 313 to allow the relative rotation. The support rod 319 assists in maintaining the radial alignment of the

[0143] Myosin is coated on a planar surface 316 of the first disc 311 that is obverse to the disc surface secured to the base 312. Actin is coated on a planar surface 317 of the second disc 313 that is obverse to the disc surface secured